

IN THE CLAIMS

This listing of claims below will replace all prior versions and listings of claims in this application:

Listing of Claims

1-4. (Canceled)

5. (currently amended): A transgenic plant transformed with a DNA that encodes a protein consisting of the amino acid sequence as shown in SEQ ID NO: 8, operably linked downstream of a stress responsive promoter comprising a DRE region.

6. (currently amended): A transgenic plant transformed with a DNA comprising the nucleotide sequence as shown in SEQ ID NO: 7, operably linked downstream of a stress responsive promoter comprising a DRE region.

7. (currently amended): A transgenic plant transformed with a DNA that encodes a protein consisting of the amino acid sequence as shown in SEQ ID NO: 8, operably linked downstream of a stress responsive promoter comprising a DRE region to which said protein can bind.

8. (currently amended) A transgenic plant transformed with a DNA, that encodes a protein, comprising the nucleotide sequence as shown in SEQ ID NO: 7 operably linked downstream of a stress responsive promoter comprising a DRE region to which said protein can bind.

9. (new): The transgenic plant of claim 5, wherein the stress responsive promoter comprises at least one promoter selected from the group consisting of rd29A gene promoter, rd17 gene promoter, cor6.6 gene promoter, cor15a gene promoter, and kin1 gene promoter.

10. (new): The transgenic plant of claim 6, wherein the stress responsive promoter comprises at least one promoter selected from the group consisting of rd29A gene promoter, rd17 gene promoter, cor6.6 gene promoter, cor15a gene promoter, and kin1 gene promoter.

11. (new): The transgenic plant of claim 7, wherein the stress responsive promoter comprises at least one promoter selected from the group consisting of rd29A gene promoter, rd17 gene promoter, cor6.6 gene promoter, cor15a gene promoter, and kin1 gene promoter.

12. (new): The transgenic plant of claim 8, wherein the stress responsive promoter comprises at least one promoter selected from the group consisting of rd29A gene promoter, rd17 gene promoter, cor6.6 gene promoter, cor15a gene promoter, and kin1 gene promoter.